RAW SEQUENCE LISTING
PATENT APPLICATION *US/08/061,699A*

INPUT SET: S2691.raw

> This Raw Listing contains the General
> Information Section and those Sequences
> containing ERRORS.

```
1                                       SEQUENCE LISTING
2
3    (1)      General Information:
4
5         (i) APPLICANT: Weiner, Amy J.
6                        Houghton, Michael
7
8       (ii) TITLE OF INVENTION: Conserved Motif of Hepatitis C Virus
9              E2/NS1 Region
10
11    (iii) NUMBER OF SEQUENCES: 133
12
13     (iv) CORRESPONDENCE ADDRESS:
14            (A) ADDRESSEE: Chiron Corporation
15            (B) STREET: 4560 Horton Street
16            (C) CITY: Emeryville
17            (D) STATE: CA
18            (E) COUNTRY: USA
19            (F) ZIP: 94608-2916
20
21      (v) COMPUTER READABLE FORM:
22            (A) MEDIUM TYPE: Floppy disk
23            (B) COMPUTER: IBM PC compatible
24            (C) OPERATING SYSTEM: PC-DOS/MS-DOS
25            (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
26
27     (vi) CURRENT APPLICATION DATA:
28            (A) APPLICATION NUMBER: US 08/061,699
29            (B) FILING DATE: 12-MAY-1993
30            (C) CLASSIFICATION:
31
32   (viii) ATTORNEY/AGENT INFORMATION:
33            (A) NAME: Goldman, Kenneth M.
34            (B) REGISTRATION NUMBER: 34,174
35            (C) REFERENCE/DOCKET NUMBER: 0938.001
36
37     (ix) TELECOMMUNICATION INFORMATION:
38            (A) TELEPHONE: 510-601-2719
39            (B) TELEFAX: 510-655-3542
40
41
```

ERRORED SEQUENCES FOLLOW:

```
 84    (2) INFORMATION FOR SEQ ID NO:2:
 85
 86        (i) SEQUENCE CHARACTERISTICS:
 87            (A) LENGTH: 31 amino acids
 88            (B) TYPE: amino acid
 89            (C) STRANDEDNESS: single
 90            (D) TOPOLOGY: linear
 91
 92        (ii) MOLECULE TYPE: peptide
 93
 94        (ix) FEATURE:
 95            (A) NAME/KEY: Ahe
 96            (B) LOCATION: 2
 97            (D) OTHER INFORMATION: /label= variable
 98                        /note= "2-amino heptanoic is butyl alanine"
 99
100
101        (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
102
103        Cys Ahe Glu Thr His Val Thr Gly Gly Ser Ala Gly His Thr Val Ser
104        1               5                   10                  15
105
106        Gly Phe Val Ser Leu Leu Ala Pro Gly Ala Lys Gln Asn Val Gln Leu
107                        20              25                  30
108
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(handwritten annotations:) Xaa — Represent this as an Xaa. "Ahe" is not valid to have in a sequence.

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION *US/08/061,699A*

INPUT SET: S2691.raw

Line	Error	Original Text
103	Wrong Amino Acid Designator	Cys Ahe Glu Thr His Val Thr Gly Gly Ser Ala Gly His Thr